

SEQUENCE LISTING

<110> AVENTIS PHARMA SA
US GOVERNMENT of the UNITED STATES

<120> NUCLEIC ACID OF THE HUMAN ABCC11 GENE, VECTORS
CONTAINING SUCH NUCLEIC ACID, AND USES THEREOF

<130> ABCC11 GENE

<140> ST01005

<141> 2002-03-05

<150> 60/272,757

<151> 2001-03-05

<160> 31

<170> PatentIn Ver. 2.1

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				580												
Leu	Met	Gly	Gly	Ala	Tyr	Asp	Lys	Ala	Arg	Tyr	Leu	Gln	Val	Leu	His	
				595					600					605		
Cys	Cys	Ser	Leu	Asn	Arg	Asp	Leu	Glu	Leu	Leu	Pro	Phe	Gly	Asp	Met	
				610					615					620		
Thr	Glu	Ile	Gly	Glu	Arg	Gly	Leu	Asn	Leu	Ser	Gly	Gly	Gln	Lys	Gln	
				625					630					635		
Arg	Ile	Ser	Leu	Ala	Arg	Ala	Val	Tyr	Ser	Asp	Arg	Gln	Ile	Tyr	Leu	
				645					650					655		
Leu	Asp	Asp	Pro	Leu	Ser	Ala	Val	Asp	Ala	His	Val	Gly	Lys	His	Ile	
				660					665					670		
Phe	Glu	Glu	Cys	Ile	Lys	Lys	Thr	Leu	Arg	Gly	Lys	Thr	Val	Val	Leu	
				675					680					685		
Val	Thr	His	Gln	Leu	Gln	Tyr	Leu	Glu	Phe	Cys	Gly	Gln	Ile	Ile	Leu	
				690					695					700		
Leu	Glu	Asn	Gly	Lys	Ile	Cys	Glu	Asn	Gly	Thr	His	Ser	Glu	Leu	Met	
				705					710					715		
Gln	Lys	Lys	Gly	Lys	Tyr	Ala	Gln	Leu	Ile	Gln	Lys	Met	His	Lys	Glu	
				725					730					735		
Ala	Thr	Ser	Asp	Met	Leu	Gln	Asp	Thr	Ala	Lys	Ile	Ala	Glu	Lys	Pro	
				740					745					750		
Lys	Val	Glu	Ser	Gln	Ala	Leu	Ala	Thr	Ser	Leu	Glu	Glu	Ser	Leu	Asn	
				755					760					765		
Gly	Asn	Ala	Val	Pro	Glu	His	Gln	Leu	Thr	Gln	Glu	Glu	Glu	Met	Glu	
				770					775					780		
Glu	Gly	Ser	Leu	Ser	Trp	Arg	Val	Tyr	His	His	Tyr	Ile	Gln	Ala	Ala	
				785					790					795		
Gly	Gly	Tyr	Met	Val	Ser	Cys	Ile	Ile	Phe	Phe	Phe	Val	Val	Leu	Ile	
				805					810					815		
Val	Phe	Leu	Thr	Ile	Phe	Ser	Phe	Trp	Trp	Leu	Ser	Tyr	Trp	Leu	Glu	
				820					825					830		
Gln	Gly	Ser	Gly	Thr	Asn	Ser	Ser	Arg	Glu	Ser	Asn	Gly	Thr	Met	Ala	
				835					840					845		
Asp	Leu	Gly	Asn	Ile	Ala	Asp	Asn	Pro	Gln	Leu	Ser	Phe	Tyr	Gln	Leu	
				850					855					860		
Val	Tyr	Gly	Leu	Asn	Ala	Leu	Leu	Leu	Ile	Cys	Val	Gly	Val	Cys	Ser	
				865					870					875		
Ser	Gly	Ile	Phe	Thr	Lys	Val	Thr	Arg	Lys	Ala	Ser	Thr	Ala	Leu	His	

885						890						895					
Asn	Lys	Leu	Phe	Asn	Lys	Val	Phe	Arg	Cys	Pro	Met	Ser	Phe	Phe	Asp		
			900					905						910			
Thr	Ile	Pro	Ile	Gly	Arg	Leu	Leu	Asn	Cys	Phe	Ala	Gly	Asp	Leu	Glu		
		915					920					925					
Gln	Leu	Asp	Gln	Leu	Leu	Pro	Ile	Phe	Ser	Glu	Gln	Phe	Leu	Val	Leu		
	930					935					940						
Ser	Leu	Met	Val	Ile	Ala	Val	Leu	Leu	Ile	Val	Ser	Val	Leu	Ser	Pro		
945					950					955					960		
Tyr	Ile	Leu	Leu	Met	Gly	Ala	Ile	Ile	Met	Val	Ile	Cys	Phe	Ile	Tyr		
				965					970					975			
Tyr	Met	Met	Phe	Lys	Lys	Ala	Ile	Gly	Val	Phe	Lys	Arg	Leu	Glu	Asn		
			980					985					990				
Tyr	Ser	Arg	Ser	Pro	Leu	Phe	Ser	His	Ile	Leu	Asn	Ser	Leu	Gln	Gly		
		995					1000					1005					
Leu	Ser	Ser	Ile	His	Val	Tyr	Gly	Lys	Thr	Glu	Asp	Phe	Ile	Ser	Gln		
	1010					1015					1020						
Phe	Lys	Arg	Leu	Thr	Asp	Ala	Gln	Asn	Asn	Tyr	Leu	Leu	Leu	Phe	Leu		
1025					1030					1035					1040		
Ser	Ser	Thr	Arg	Trp	Met	Ala	Leu	Arg	Leu	Glu	Ile	Met	Thr	Asn	Leu		
				1045					1050					1055			
Val	Thr	Leu	Ala	Val	Ala	Leu	Phe	Val	Ala	Phe	Gly	Ile	Ser	Ser	Thr		
		1060						1065					1070				
Pro	Tyr	Ser	Phe	Lys	Val	Met	Ala	Val	Asn	Ile	Val	Leu	Gln	Leu	Ala		
	1075						1080					1085					
Ser	Ser	Phe	Gln	Ala	Thr	Ala	Arg	Ile	Gly	Leu	Glu	Thr	Glu	Ala	Gln		
	1090					1095					1100						
Phe	Thr	Ala	Val	Glu	Arg	Ile	Leu	Gln	Tyr	Met	Lys	Met	Cys	Val	Ser		
1105					1110					1115					1120		
Glu	Ala	Pro	Leu	His	Met	Glu	Gly	Thr	Ser	Cys	Pro	Gln	Gly	Trp	Pro		
				1125					1130					1135			
Gln	His	Gly	Glu	Ile	Ile	Phe	Gln	Asp	Tyr	His	Met	Lys	Tyr	Arg	Asp		
		1140						1145					1150				
Asn	Thr	Pro	Thr	Val	Leu	His	Gly	Ile	Asn	Leu	Thr	Ile	Arg	Gly	His		
		1155					1160					1165					
Glu	Val	Val	Gly	Ile	Val	Gly	Arg	Thr	Gly	Ser	Gly	Lys	Ser	Ser	Leu		
	1170					1175					1180						
Gly	Met	Ala	Leu	Phe	Arg	Leu	Val	Glu	Pro	Met	Ala	Gly	Arg	Ile	Leu		
1185					1190					1195					1200		
Ile	Asp	Gly	Val	Asp	Ile	Cys	Ser	Ile	Gly	Leu	Glu	Asp	Leu	Arg	Ser		
				1205					1210					1215			

Lys Leu Ser Val Ile Pro Gln Asp Pro Val Leu Leu Ser Gly Thr Ile
 1220 1225 1230
 Arg Phe Asn Leu Asp Pro Phe Asp Arg His Thr Asp Gln Gln Ile Trp
 1235 1240 1245
 Asp Ala Leu Glu Arg Thr Phe Leu Thr Lys Ala Ile Ser Lys Phe Pro
 1250 1255 1260
 Lys Lys Leu His Thr Asp Val Val Glu Asn Gly Gly Asn Phe Ser Val
 1265 1270 1275 1280
 Gly Glu Arg Gln Leu Leu Cys Ile Ala Arg Ala Val Leu Arg Asn Ser
 1285 1290 1295
 Lys Ile Ile Leu Ile Asp Glu Ala Thr Ala Ser Ile Asp Met Glu Thr
 1300 1305 1310
 Asp Thr Leu Ile Gln Arg Thr Ile Arg Glu Ala Phe Gln Gly Cys Thr
 1315 1320 1325
 Val Leu Val Ile Ala His Arg Val Thr Thr Val Leu Asn Cys Asp His
 1330 1335 1340
 Ile Leu Val Met Gly Asn Gly Lys Val Val Glu Phe Asp Arg Pro Glu
 1345 1350 1355 1360
 Val Leu Arg Lys Lys Pro Gly Ser Leu Phe Ala Ala Leu Met Ala Thr
 1365 1370 1375
 Ala Thr Ser Ser Leu Arg
 1380